IOTECHNOLOGY
SYSTEMS
BRANCH

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

Source

Date Processed by STIC:

10/823,964 TEWD

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221 Effective 12/13/03: TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER VERSION 4.1 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 2023 1 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission & User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22312-1459
- Hand Carry directly to (EFFECTIVE 12/0)/03):
 U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two. 2011 South Clark Place, Arlington, VA 22202
- 4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1803-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03

BEST AVAILABLE COPY

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION SERIAL NUMBER: 10/823,964
ATTN: NEW RULES CASES	: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARD
lWrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.
3Misaligned Amino Numbering	The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
4Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
5Variable Length	Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each nor Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6Patentin 2.0 "bug"	A "bug" in Patentin version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s)
7Skipped Sequences (OLD RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence. (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped
	Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences
8Skipped Sequences (NEW RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence <210> sequence id number <400> sequence id number
	000
9Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
10Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
11 Use of <220>	Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 00/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
Patentin 2.0 "bug"	Please do not use "Copy to Disk" function of Patentln version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
13 Misuse of n/Xaa	"n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid
	AMC - Biotechnology Systems Branch - 09/09/2003



EF

RAW SEQUENCE LISTING DATE: 04/21/2004 PATENT APPLICATION: US/10/823,964 TIME: 07:41:49

Input Set : A:\Seqlist.txt

4 <110> APPLICANT: BAM, NARENDRA

Output Set: N:\CRF4\04212004\J823964.raw

```
BONGERS, JACOB
6
        KIRKPATRICK, ROBERT B.
7
        JANSON, CHERYL A.
8
        JOHANSON, KYUNG
        QIU, XIANYANG
        YEH, PING
  <120> TITLE OF INVENTION: CONJUGATES COMPRISING HUMAN IL-18 AND
: 3
        SUBSTITUTION MUTANTS THEREOF
16 <130> FILE REFERENCE: PU60053
18 <140> CURRENT APPLICATION NUMBER: US/10/823,964
19 <141> CURRENT FILING DATE: 2004-04-14
11 <150> PRIOR APPLICATION NUMBER: 60/462,947
                                                         Does Not Comply
22 <151> PRIOR FILING DATE: 2003-04-15
                                                         Corrected Diskette Needed
24 <160> NUMBER OF SEQ ID NOS: 28
26 <170> SOFTWARE: FastSEQ for Windows Version 4.0
28 <210> SEQ ID NO: 1
19 <211> LENGTH: 157
30 <212> TYPE: PRT
31 <213> ORGANISM: Homo sapiens
33 <400> SEQUENCE: 1
34 Tyr Phe Gly Lys Leu Glu Ser Lys Leu Ser Val Ile Arg Asn Leu Asn
35 1
                 5
36 Asp Gln Val Leu Phe Ile Asp Gln Gly Asn Arg Pro Leu Phe Glu Asp
37
             20
                                 25
38 Met Thr Asp Ser Asp Cys Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile
  35
39
                             40
10 Ile Ser Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile
11
                                            60
12 Ser Val Lys Cys Glu Lys Ile Ser Thr Leu Ser Cys Glu Asn Lys Ile
13 65
                    70
                                     75
14 Ile Ser Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp Thr Lys
15
                  85
                                     90
16 Ser Asp Ile Ile Phe Phe Glm Arg Ser Val Pro Gly His Asp Asm Lys
17
                                 105
48 Met Gln Phe Glu Ser Ser Ser Tyr Glu Gly Tyr Phe Leu Ala Cys Glu
19 115
                             120
50 Lys Glu Arg Asp Leu Phe Lys Leu Ile Leu Lys Lys Glu Asp Glu Leu
51 130
                        135.
32 Gly Asp Arg Ser Ile Met Phe Thr Val Gln Asn Glu Asp
33 145
                     150
36 <210> SEQ ID NO: 2
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57 <211> LENGTH: 157

RAW SEQUENCE LISTING DATE: 04/21/2004 PATENT APPLICATION: US/10/823,964 TIME: 07:41:49

Input Set : A:\Seqlist.txt

Output Set: N:\CRF4\04212004\J823964.raw

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58 <212> TYPE: PRT
59 <213> ORGANISM: Mus musculus
51 <400> SEQUENCE: 2
52 Asn Phe Gly Arg Leu His Cys Thr Thr Ala Val Ile Arg Asn Ile Asn
53 1
                                       10
54 Asp Gln Val Leu Phe Val Asp Lys Arg Gln Pro Val Phe Glu Asp Met
55
              20
56 Thr Asp Ile Asp Gln Ser Ala Ser Glu Pro Gln Thr Arg Leu Ile Ile
          35
57
                               40
  Tyr Met Tyr Lys Asp Ser Glu Val Arg Gly Leu Ala Val Thr Leu Ser
59
                          55
70 Val Lys Asp Ser Lys Met Ser Thr Leu Ser Cys Lys Asn Lys Ile Ile
71 65
72 Ser Phe Glu Glu Met Asp Pro Pro Glu Asn Ile Asp Asp Ile Gln Ser
73
74 Asp Leu Ile Phe Phe Gln Lys Arg Val Pro Gly His Asn Lys Met Glu
75
                                   105
                                                       110
              100
76 Phe Glu Ser Ser Leu Tyr Glu Gly His Phe Leu Ala Cys Gln Lys Glu
77
         115
                               120
                                                   125
78 Asp Asp Ala Phe Lys Leu Ile Leu Lys Lys Lys Asp Glu Asn Gly Asp
79
     130
                           135
                                               140
30 Lys Ser Val Met Phe Thr Leu Thr Asn Leu His Gln Ser
81 145
                       150
84 <210> SEQ ID NO: 3
85 <211> LENGTH: 203
86 <212> TYPE: PRT
87 <213> ORGANISM: Homo sapiens
89 <400> SEQUENCE: 3
90 Met His His His His His Thr Arg Gly Met Ala Ala Glu Pro Val
92 Glu Asp Asn Cys Ile Asn Phe Val Ala Met Lys Phe Ile Asp Asn Thr
93
              20
                                   25
94 Leu Tyr Phe Ile Ala Glu Asp Asp Glu Asn Leu Glu Ser Asp Tyr Phe
           35
96 Gly Lys Leu Glu Ser Lys Leu Ser Val Ile Arg Asn Leu Asn Asp Gln
       50
                          55
98 Val Leu Phe Ile Asp Gln Gly Asn Arg Pro Leu Phe Glu Asp Met Thr
                       70
                                           75
100 Asp Ser Asp Cys Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile Ile Ser
101
                    85
102 Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile Ser Val
103
                100
                                    105
104 Lys Cys Glu Lys Ile Ser Thr Leu Ser Cys Glu Asn Lys Ile Ile Ser
                                120
           115
106 Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp Thr Lys Ser Asp
      130
                            135
                                                140
108 Ile Ile Phe Phe Gln Arg Ser Val Pro Gly His Asp Asn Lys Met Gln
```

150

110 Phe Glu Ser Ser Tyr Glu Gly Tyr Phe Leu Ala Cys Glu Lys Glu

155

109 145

PATENT APPLICATION: US/10/823,964 TIME: 07:41:49 Input Set : A:\Seqlist.txt Output Set: N:\CRF4\04212004\J823964.raw 165 tem errormary
human Sheet. 111 170 112 Arg Asp Leu Phe Lys Leu Ile Leu Lys Lys Glu Asp Glu Leu Gly Asp 185 114 Arg Ser Ile Met Phe Thr Val Gln Asn Glu Asp 195 please explain. 115 200 118 <210> SEQ ID NO: 4 119 <211> LENGTH: 157 120 <212> TYPE: PRT 121 <213> ORGANISM: (Artificial Sequence 123 <220> FEATURE: 124 <223> OTHER INFORMATION: Whereby the Cysteine at position 38 of the human 125 IL-18 sequence has been replaced with Serine. 127 <400> SEQUENCE: 4 128 Tyr Phe Gly Lys Leu Glu Ser Lys Leu Ser Val Ile Arg Asn Leu Asn 129 1 130 Asp Gln Val Leu Phe Ile Asp Gln Gly Asn Arg Pro Leu Phe Glu Asp 132 Met Thr Asp Ser Asp Ser Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile 35 133 40 45 134 Ile Ser Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile 136 Ser Val Lys Cys Glu Lys Ile Ser Thr Leu Ser Cys Glu Asn Lys Ile 70 75 138 Ile Ser Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp Thr Lys 85 90 140 Ser Asp Ile Ile Phe Phe Gln Arg Ser Val Pro Gly His Asp Asn Lys 141 100 105 Per New Sequence mules, Po For New Sequence maximum (2232)

ine at position 38 of the human aced with position 2000. 110 142 Met Gln Phe Glu Ser Ser Tyr Glu Gly Tyr Phe Leu Ala Cys Glu 143 115 120 144 Lys Glu Arg Asp Leu Phe Lys Leu Ile Leu Lys Lys Glu Asp Glu Leu 135 146 Gly Asp Arg Ser Ile Met Phe Thr Val Gln Asn Glu Asp 147 145 150 150 <210> SEQ ID NO: 5 151 <211> LENGTH: 157 152 <212> TYPE: PRT 153 <213> ORGANISM: Artificial Sequence 155 <220> FEATURE: 156 <223> OTHER INFORMATION: Whereby the Cysteine at position 38 of the human 157 11-18 sequence has been replaced with Serine, the

Cysteine at position 68 has been replaced with,

163 Tyr Phe Gly Lys Leu Glu Ser Lys Leu Ser Val Ile Arg Asn Leu Asn

165 Asp Gln Val Leu Phe Ile Asp Gln Gly Asn Arg Pro Leu Phe Glu Asp

167 Met Thr Asp Ser Asp Ser Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile

25

Aspartic acid, and the Asparagine at position (78 220) has been replaced with Cysteine.

5

20

DATE: 04/21/2004

RAW SEQUENCE LISTING

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DATE: 04/21/2004

TIME: 07:41:49

```
Input Set : A:\Seqlist.txt
                                                                       7 se p5.3
                Output Set: N:\CRF4\04212004\J823964.raw
   Ile Ser Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile
171 Ser Val Lys Asp Glu Lys Ile Ser Thr Leu Ser Cys Glu (Asn) Lys Ile
172 65
                        70
173 Ile Ser Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp Thr Lys
174
                                         90
175 Ser Asp Ile Ile Phe Phe Gln Arg Ser Val Pro Gly His Asp Asn Lys
176
               100
                                    105
177 Met Gln Phe Glu Ser Ser Ser Tyr Glu Gly Tyr Phe Leu Ala Cys Glu
L78
           115
                                120
                                                     125
179 Lys Glu Arg Asp Leu Phe Lys Leu Ile Leu Lys Lys Glu Asp Glu Leu
180
                            135
                                                 140
181 Gly Asp Arg Ser Ile Met Phe Thr Val Gln Asn Glu Asp
                        150
185 <210> SEQ ID NO: 6
186 <211> LENGTH: 157
187 <212> TYPE: PRT
188 <213> ORGANISM: Artificial Sequence
190 <220> FEATURE:
191 <223 TOTHER INFORMATION: Whereby the Cysteine at position 38 of the human
         ) IL-18 sequence has been replaced with Serine, the
192
        Cysteine at position 68 has been replaced with
193
    Aspartic acid, and the Glutamic acid at position
195 233121 has been 197 <400> SEQUENCE: 6
      23)121 has been replaced with Cysteine.
198 Tyr Phe Gly Lys Leu Glu Ser Lys Leu Ser Val Ile Arg Asn Leu Asn
199 1
200 Asp Gln Val Leu Phe Ile Asp Gln Gly Asn Arg Pro Leu Phe Glu Asp
               20
                                    25
202 Met Thr Asp Ser Asp Ser Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile
203
           35
                                40
204 Ile Ser Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile
                            55
206 Ser Val Lys Asp Glu Lys Ile Ser Thr Leu Ser Cys Glu Asn Lys Ile
207 65
                        70
108 Ile Ser Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp Thr Lys
                    85
                                         90
210 Ser Asp Ile Ile Phe Phe Gln Arg Ser Val Pro Gly His Asp Asn Lys
211
               100
                                    105
                                                         110
212 Met Gln Phe Glu Ser Ser Ser Tyr Cys Gly Tyr Phe Leu Ala Cys Glu
           115
                                120
                                                     125
214 Lys Glu Arg Asp Leu Phe Lys Leu Ile Leu Lys Lys Glu Asp Glu Leu
       130
                            135
                                                 140
216 Gly Asp Arg Ser Ile Met Phe Thr Val Gln Asn Glu Asp
217 145
220 <210> SEQ ID NO: 7
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US/10/823,964

RAW SEQUENCE LISTING

PATENT APPLICATION:

221 <211> LENGTH: 157 222 <212> TYPE: PRT

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Input Set : A:\Seqlist.txt
                Output Set: N:\CRF4\04212004\J823964.raw
223 <213> ORGANISM: Artificial Sequence
?25 <220> FEATURE:
226 <223> OTHER INFORMATION: Whereby the Cysteine at position 38 of the human
227
          IL-18 sequence has been replaced with Serine, the
228
          Cysteine at position 68 has been replaced with
    Aspartic acid, and the Leucine at position 144 has Some error
230 (253) been replaced with Cysteine.
232 <400> SEQUENCE: 7
233 Tyr Phe Gly Lys Leu Glu Ser Lys Leu Ser Val Ile Arg Asn Leu Asn
                                     . ... 10
235 Asp Gln Val Leu Phe Ile Asp Gln Gly Asn Arg Pro Leu Phe Glu Asp
336
                20
                                    25
237
   Met Thr Asp Ser Asp Ser Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile
238
            35
   Ile Ser Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile
341 Ser Val Lys Asp Glu Lys Ile Ser Thr Leu Ser Cys Glu Asn Lys Ile
                        70
143 Ile Ser Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp Thr Lys
244
                    85
                                         90
:45 Ser Asp Ile Ile Phe Phe Gln Arg Ser Val Pro Gly His Asp Asn Lys
246
                100
                                    105
247 Met Gln Phe Glu Ser Ser Ser Tyr Glu Gly Tyr Phe Leu Ala Cys Glu
            115
                                120
249 Lys Glu Arg Asp Leu Phe Lys Leu Ile Leu Lys Lys Glu Asp Glu Cys
250
       130
                            135
351 Gly Asp Arg Ser Ile Met Phe Thr Val Gln Asn Glu Asp
252 145
255 <210> SEQ ID NO: 8
256 <211> LENGTH: 157
257 <212> TYPE: PRT
258 <213> ORGANISM: Artificial Sequence
260 <220> FEATURE:
261 <223> OTHER INFORMATION: Whereby the Cysteine at position 38 of the human
          IL-18 sequence has been replaced with Serine, the
262
263
          Cysteine at position 68 has been replaced with
Aspartic acid, Aspartic acid at position 157 has been replaced with Cysteine.
267 <400 SEQUENCE: 8
268 Tyr Phe Gly Lys Leu Glu Ser Lys Leu Ser Val Ile Arg Asn Leu Asn
269
   1
                     5
                                        10
270 Asp Gln Val Leu Phe Ile Asp Gln Gly Asn Arg Pro Leu Phe Glu Asp
                20
                                    25
   Met Thr Asp Ser Asp Ser Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile
                                40
   Ile Ser Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile
       50
                            55
276 Ser Val Lys Asp Glu Lys Ile Ser Thr Leu Ser Cys Glu Asn Lys Ile
277 65
```

The type of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

DATE: 04/21/2004

TIME: 07:41:49

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/823,964

:\CRF4\Outhold\VsrJ823964.htm

4/21,

VERIFICATION SUMMARY DATE: 04/21/2004
PATENT APPLICATION: US/10/823,964 TIME: 07:41:50

Input Set : A:\Seglist.txt

Output Set: N:\CRF4\04212004\J823964.raw

M:270 C: Current Application Number differs, Replaced Current Application Number M:259 W: Allowed number of lines exceeded, <223> Other Information:
M:259 W: Allowed number of lines exceeded, <223> Other Information:
M:259 W: Allowed number of lines exceeded, <223> Other Information:
M:259 W: Allowed number of lines exceeded, <223> Other Information:
M:259 W: Allowed number of lines exceeded, <223> Other Information:
M:259 W: Allowed number of lines exceeded, <223> Other Information: